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SEQUENCE LISTING

<110> ULLRICH, Axel
NAYLER, Oliver

<120> CLK PROTEIN KINASES AND RELATED PRODUCTS AND METHODS

<130> 038602/0431

<140> US 09/127,248

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<160> 26

<170> PatentIn version 3.0

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39

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39

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<400> 12
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36

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24

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<400> 16
gttgccctga ggatcatccg gaat

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 <223> Xaa at position 2 can be Val or Met

<220>
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 <211> 483
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 <213> Mus musculus

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 20 25 30

Arg Ser His Ser Ser Ala Arg Glu Gln Lys Arg Cys Arg Tyr Asp His
 35 40 45
 Ser Lys Thr Thr Asp Ser Tyr Tyr Leu Glu Ser Arg Ser Ile Asn Glu
 50 55 60
 Lys Ala Tyr His Ser Arg Arg Tyr Val Asp Glu Tyr Arg Asn Asp Tyr
 65 70 75 80
 Met Gly Tyr Glu Pro Gly His Pro Tyr Gly Glu Pro Gly Ser Arg Tyr
 85 90 95
 Gln Met His Ser Ser Lys Ser Ser Gly Arg Ser Gly Arg Ser Ser Tyr
 100 105 110
 Lys Ser Lys His Arg Ser Arg His His Thr Ser Gln His His Ser His
 115 120 125
 Gly Lys Ser His Arg Arg Lys Arg Ser Arg Ser Val Glu Asp Asp Glu
 130 135 140
 Glu Gly His Leu Ile Cys Gln Ser Gly Asp Val Leu Ser Ala Arg Tyr
 145 150 155 160
 Glu Ile Val Asp Thr Leu Gly Glu Gly Ala Phe Gly Lys Val Val Glu
 165 170 175
 Cys Ile Asp His Lys Val Gly Gly Arg Arg Val Ala Val Lys Ile Val
 180 185 190
 Lys Asn Val Asp Arg Tyr Cys Glu Ala Ala Gln Ser Glu Ile Gln Val
 195 200 205
 Leu Glu His Leu Asn Thr Thr Asp Pro His Ser Thr Phe Arg Cys Val
 210 215 220
 Gln Met Leu Glu Trp Phe Glu His Arg Gly His Ile Cys Ile Val Phe
 225 230 235 240
 Glu Leu Leu Gly Leu Ser Thr Tyr Asp Phe Ile Lys Glu Asn Ser Phe
 245 250 255
 Leu Pro Phe Arg Met Asp His Ile Arg Lys Met Ala Tyr Gln Ile Cys
 260 265 270
 Lys Ser Val Asn Phe Leu His Ser Met Lys Leu Thr His Thr Asp Leu
 275 280 285
 Lys Pro Glu Asn Ile Leu Phe Val Lys Ser Asp Tyr Thr Glu Ala Tyr
 290 295 300
 Asn Pro Lys Met Lys Arg Asp Glu Arg Thr Ile Val Asn Pro Asp Ile
 305 310 315 320

Lys Val Val Asp Phe Gly Ser Ala Thr Tyr Asp Asp Glu His His Ser
 325 330 335
 Thr Leu Val Ser Thr Arg His Tyr Arg Ala Pro Glu Val Ile Leu Ala
 340 345 350
 Leu Gly Trp Ser Gln Pro Cys Asp Val Trp Ser Ile Gly Cys Ile Leu
 355 360 365
 Ile Glu Tyr Tyr Leu Gly Phe Thr Val Phe Pro Thr His Asp Ser Arg
 370 375 380
 Glu His Leu Ala Met Met Glu Arg Ile Leu Gly Pro Leu Pro Lys His
 385 390 395 400
 Met Ile Gln Lys Thr Arg Lys Arg Arg Tyr Phe His His Asp Arg Leu
 405 410 415
 Asp Trp Asp Glu His Ser Ser Ala Gly Arg Tyr Val Ser Arg Arg Cys
 420 425 430
 Lys Pro Leu Lys Glu Phe Met Leu Ser Gln Asp Ala Glu His Glu Phe
 435 440 445
 Leu Phe Asp Leu Val Gly Lys Ile Leu Glu Tyr Asp Pro Ala Lys Arg
 450 455 460
 Ile Thr Leu Lys Glu Ala Leu Lys His Pro Phe Phe Tyr Pro Leu Lys
 465 470 475 480
 Lys His Thr

<210> 21
 <211> 499
 <212> PRT
 <213> Mus musculus

<400> 21

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 20 25 30
 Arg Ser Trp Ser Ser Ser Ser Asp Arg Thr Arg Arg Arg Arg Glu
 35 40 45
 Asp Ser Tyr His Val Arg Ser Arg Ser Ser Tyr Asp Asp His Ser Ser
 50 55 60
 Asp Arg Arg Leu Tyr Asp Arg Arg Tyr Cys Gly Ser Tyr Arg Arg Asn
 65 70 75 80

Asp Tyr Ser Arg Asp Arg Gly Glu Ala Tyr Tyr Asp Thr Asp Phe Arg
 85 90 95
 Gln Ser Tyr Glu Tyr His Arg Glu Asn Ser Ser Tyr Arg Ser Gln Arg
 100 105 110
 Ser Ser Arg Arg Lys His Arg Arg Arg Arg Arg Ser Arg Thr Phe
 115 120 125
 Ser Arg Ser Ser Ser His Ser Ser Arg Arg Ala Lys Ser Val Glu Asp
 130 135 140
 Asp Ala Glu Gly His Leu Ile Tyr His Val Gly Asp Trp Leu Gln Glu
 145 150 155 160
 Arg Tyr Glu Ile Val Ser Thr Leu Gly Glu Gly Thr Ser Gly Arg Val
 165 170 175
 Val Gln Cys Val Asp His Arg Arg Gly Gly Thr Arg Val Ala Leu Lys
 180 185 190
 Ile Ile Lys Asn Val Glu Lys Tyr Lys Glu Ala Ala Arg Leu Glu Ile
 195 200 205
 Asn Val Leu Glu Lys Ile Asn Glu Lys Asp Pro Asp Asn Lys Asn Leu
 210 215 220
 Cys Val Gln Met Phe Asp Trp Phe Asp Tyr His Gly His Met Cys Ile
 225 230 235 240
 Ser Phe Glu Leu Leu Gly Leu Ser Thr Phe Asp Phe Leu Lys Asp Asn
 245 250 255
 Asn Tyr Leu Pro Tyr Pro Ile His Gln Val Arg His Met Ala Phe Gln
 260 265 270
 Leu Cys Gln Ala Val Lys Phe Leu His Asp Asn Lys Leu Thr His Thr
 275 280 285
 Asp Leu Lys Pro Glu Asn Ile Leu Phe Val Asn Ser Asp Tyr Glu Leu
 290 295 300
 Thr Tyr Asn Leu Glu Lys Lys Arg Asp Glu Arg Ser Val Lys Ser Thr
 305 310 315 320
 Ala Val Arg Val Val Asp Phe Gly Ser Ala Thr Phe Asp His Glu His
 325 330 335
 His Ser Thr Ile Val Ser Thr Arg His Tyr Arg Ala Pro Glu Val Ile
 340 345 350
 Leu Glu Leu Gly Trp Ser Gln Pro Cys Asp Val Trp Ser Ile Gly Cys
 355 360 365

Ile Ile Phe Glu Tyr Tyr Val Gly Phe Thr Leu Phe Gln Thr His Asp
 370 375 380
 Asn Arg Glu His Leu Ala Met Met Glu Arg Ile Leu Gly Pro Val Pro
 385 390 395 400
 Ser Arg Met Ile Arg Lys Thr Arg Lys Gln Lys Tyr Phe Tyr Arg Gly
 405 410 415
 Arg Leu Asp Trp Asp Glu Asn Thr Ser Ala Gly Arg Tyr Val Arg Glu
 420 425 430
 Asn Cys Lys Pro Leu Arg Arg Tyr Leu Thr Ser Glu Ala Glu Asp His
 435 440 445
 His Gln Leu Phe Asp Leu Ile Glu Asn Met Leu Glu Tyr Glu Pro Ala
 450 455 460
 Lys Arg Leu Thr Leu Gly Glu Ala Leu Gln His Pro Phe Phe Ala Cys
 465 470 475 480
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 Ile Ser Arg

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 <212> DNA
 <213> Mus musculus

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 cgctcctggt caagtagcag tgaccggaca aggcggcggc ggagggagga cagctaccac 180
 gttcgggtccc gaagcagcta tgatgaccat tcgtccgac gccgggctgta cgatcggcgg 240
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 acagactttc ggcagtccta tgaataccat cgagagaaca gcagttaccg aagccagcgc 360
 agcagccgaa ggaaacacag aaggcggagg agacggagcc ggacattcag ccgctcatct 420
 tcacacagca gccggagagc caagagtgtg gaggacgacg ctgagggcca cctcatctac 480
 cagtcggggg actgggtaca agagcgatat gaaattgtaa gcaccttagg agaagggact 540
 tcgggcccag ttgtgcagtg tgtggaccat cgcaggggcg gaacacgagt tgccctgaag 600
 atcattaaga atgtggagaa gtacaaggaa gcagcccgc tagaaatcaa cgtgctggag 660

aaaatcaatg agaaagatcc tgacaacaag aacctctgtg tccagatggt tgactgggttt 720
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 gaaaatattc tgtttgtgaa ttcagactac gaactcacct acaacctaga gaagaagcga 960
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 ctggagttgg gctggtcaca gccatgcgat gtatggagca taggctgcat catctttgag 1140
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 gaaaggatcc tgggtcctgt cccttctcgg atgatacagaa agacaagaaa acagaaatat 1260
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 aactgcaaac ctctgcggcg gtatctgacc tcagaggcag aggaccacca ccagctcttc 1380
 gatctgattg aaaatatgct agagtatgag cctgctaagc ggctgacctt aggtgaagcc 1440
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 <211> 490
 <212> PRT
 <213> Mus musculus

<400> 23

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 Arg Leu Arg Tyr Pro Ser Arg Arg Glu Pro Pro Pro Arg Arg Ser Arg
 35 40 45
 Ser Arg Ser His Asp Arg Ile Pro Tyr Gln Arg Arg Tyr Arg Glu His
 50 55 60
 Arg Asp Ser Asp Thr Tyr Arg Cys Glu Glu Arg Ser Pro Ser Phe Gly
 65 70 75 80

Glu Asp Cys Tyr Gly Ser Ser Arg Ser Arg His Arg Arg Arg Ser Arg
 85 90 95
 Glu Arg Ala Pro Tyr Arg Thr Arg Lys His Ala His His Cys His Lys
 100 105 110
 Arg Arg Thr Arg Ser Cys Ser Ser Ala Ser Ser Arg Ser Gln Gln Ser
 115 120 125
 Ser Lys Arg Ser Ser Arg Ser Val Glu Asp Asp Lys Glu Gly His Leu
 130 135 140
 Val Cys Arg Ile Gly Asp Trp Leu Gln Glu Arg Tyr Glu Ile Val Gly
 145 150 155 160
 Asn Leu Gly Glu Gly Thr Phe Gly Lys Val Val Glu Cys Leu Asp His
 165 170 175
 Ala Arg Gly Lys Ser Gln Val Ala Leu Lys Ile Ile Arg Asn Val Gly
 180 185 190
 Lys Tyr Arg Glu Ala Ala Arg Leu Glu Ile Asn Val Leu Lys Lys Ile
 195 200 205
 Lys Glu Lys Asp Lys Glu Asn Lys Phe Leu Cys Val Leu Met Ser Asp
 210 215 220
 Trp Phe Asn Phe His Gly His Met Cys Ile Ala Phe Glu Leu Leu Gly
 225 230 235 240
 Lys Asn Thr Phe Glu Phe Leu Lys Glu Asn Asn Phe Gln Pro Tyr Pro
 245 250 255
 Leu Pro His Val Arg His Met Ala Tyr Gln Leu Cys His Ala Leu Arg
 260 265 270
 Phe Leu His Glu Asn Gln Leu Thr His Thr Asp Leu Lys Pro Glu Asn
 275 280 285
 Ile Leu Phe Val Asn Ser Glu Phe Glu Thr Leu Tyr Asn Glu His Lys
 290 295 300
 Ser Cys Glu Glu Lys Ser Val Lys Asn Thr Ser Ile Arg Val Ala Asp
 305 310 315 320
 Phe Gly Ser Ala Thr Phe Asp His Glu His His Thr Thr Ile Val Ala
 325 330 335
 Thr Arg His Tyr Arg Pro Pro Glu Val Ile Leu Glu Leu Gly Trp Ala
 340 345 350
 Gln Pro Cys Asp Val Trp Ser Ile Gly Cys Ile Leu Phe Glu Tyr Tyr
 355 360 365

Arg Gly Phe Thr Leu Phe Gln Thr His Glu Asn Arg Glu His Leu Val
 370 375 380
 Met Met Glu Lys Ile Leu Gly Pro Ile Pro Ser His Met Ile His Arg
 385 390 395 400
 Thr Arg Lys Gln Lys Tyr Phe Tyr Lys Gly Gly Leu Val Trp Asp Glu
 405 410 415
 Asn Ser Ser Asp Gly Arg Tyr Val Lys Glu Asn Cys Lys Pro Leu Lys
 420 425 430
 Ser Tyr Met Leu Gln Asp Ser Leu Glu His Val Gln Leu Phe Asp Leu
 435 440 445
 Met Arg Arg Met Leu Glu Phe Asp Pro Ala Gln Arg Ile Thr Leu Ala
 450 455 460
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 Ser Phe His Ser Ser Arg Asn Pro Ser Arg
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 <212> DNA
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aatgtggtgc ccggtgcca gaaagcacag attggacca agcttttata tgttgtaaag 1740
ttataataaa gtgttcttac tgtttgtaaa aaaaaaaaaa aaaaaaa 1787

<210> 25
<211> 481
<212> PRT
<213> Mus musculus

<400> 25

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Ser His Ser Ser Thr Gln Glu Asn Arg His Cys Lys Pro His His Gln
35 40 45

Phe Lys Asp Ser Asp Cys His Tyr Leu Glu Ala Arg Cys Leu Asn Glu
50 55 60
Arg Asp Tyr Arg Asp Arg Arg Tyr Ile Asp Glu Tyr Arg Asn Asp Tyr
65 70 75 80
Cys Glu Gly Tyr Val Pro Arg His Tyr His Arg Asp Val Glu Ser Thr
85 90 95
Tyr Arg Ile His Cys Ser Lys Ser Ser Val Arg Ser Arg Arg Ser Ser
100 105 110
Pro Lys Arg Lys Arg Asn Arg Pro Cys Ala Ser His Gln Ser His Ser
115 120 125
Lys Ser His Arg Arg Lys Arg Ser Arg Ser Ile Glu Asp Asp Glu Glu
130 135 140
Gly His Leu Ile Cys Gln Ser Gly Asp Val Leu Arg Ala Arg Tyr Glu
145 150 155 160
Ile Val Asp Thr Leu Gly Glu Gly Ala Phe Gly Lys Val Val Glu Cys
165 170 175
Ile Asp His Gly Met Asp Gly Leu His Val Ala Val Lys Ile Val Lys
180 185 190
Asn Val Gly Arg Tyr Arg Glu Ala Ala Arg Ser Glu Ile Gln Val Leu
195 200 205
Glu His Leu Asn Ser Thr Asp Pro Asn Ser Val Phe Arg Cys Val Gln
210 215 220
Met Leu Glu Trp Phe Asp His His Gly His Val Cys Ile Val Phe Glu
225 230 235 240
Leu Leu Gly Leu Ser Thr Tyr Asp Phe Ile Lys Glu Asn Ser Phe Leu
245 250 255
Pro Phe Gln Ile Asp His Ile Arg Gln Met Ala Tyr Gln Ile Cys Gln
260 265 270
Ser Ile Asn Phe Leu His His Asn Lys Leu Thr His Thr Asp Leu Lys
275 280 285
Pro Glu Asn Ile Leu Phe Val Lys Ser Asp Tyr Val Val Lys Tyr Asn
290 295 300
Ser Lys Met Lys Arg Asp Glu Arg Thr Leu Lys Asn Thr Asp Ile Lys
305 310 315 320
Val Val Asp Phe Gly Ser Ala Thr Tyr Asp Asp Glu His His Ser Thr
325 330 335

Leu Val Ser Thr Arg His Tyr Arg Ala Pro Glu Val Ile Leu Ala Leu
 340 345 350
 Gly Trp Ser Gln Pro Cys Asp Val Trp Ser Ile Gly Cys Ile Leu Ile
 355 360 365
 Glu Tyr Tyr Leu Gly Phe Thr Val Phe Gln Thr His Asp Ser Lys Glu
 370 375 380
 His Leu Ala Met Met Glu Arg Ile Leu Gly Pro Ile Pro Ala His Met
 385 390 395 400
 Ile Gln Lys Thr Arg Lys Arg Lys Tyr Phe His His Asn Gln Leu Asp
 405 410 415
 Trp Asp Glu His Ser Ser Ala Gly Arg Tyr Val Arg Arg Arg Cys Lys
 420 425 430
 Pro Leu Lys Glu Phe Met Leu Cys His Asp Glu Glu His Glu Lys Leu
 435 440 445
 Phe Asp Leu Val Arg Arg Met Leu Glu Tyr Asp Pro Ala Arg Arg Ile
 450 455 460
 Thr Leu Asp Glu Ala Leu Gln His Pro Phe Phe Asp Leu Leu Lys Arg
 465 470 475 480
 Lys

<210> 26
 <211> 1549
 <212> DNA
 <213> Mus musculus

<400> 26
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 cactatttag aagcaagatg cttgaatgag agagattatc gggaccggag atacattgat 300
 gaatacagaa atgactactg cgaaggatat gttccaagac attaccatag agacgttgaa 360
 agcacttacc ggatccattg cagtaaattc tcagtcagga gcaggagaag cagccctaag 420
 agaaagcgta atagaccctg tgcaagtcac cagtcgcatt cgaagagcca ccgaaggaaa 480
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ctaagagcaa gatatgaaat cgtggacact ttaggtgaag gagcctttgg caaagttgta	600
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